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(e) comparing the amount of binding in the first binding mixture with the amount of binding in the second binding mixture;

wherein the compound is capable of inhibiting IL-13 binding to the IL-13 receptor when a decrease in the amount of binding of the second binding mixture occurs, and wherein said receptor chain protein comprises an amino acid sequence selected from the group consisting of:

- (1) the amino acid sequence of SEQ ID NO: 2;
- (2) the amino acid sequence of SEQ ID NO: 2 from amino acids 26 to 341; and
- (3) the amino acid sequence of SEQ ID NO: 2 from amino acids 363 to 380.
- 61. (New) The method of claim 60, wherein said protein comprises the amino acid sequence of SEQ ID NO: 2.
- 62. (New) The method of claim 60, wherein said protein comprises the amino acid sequence of SEO ID NO: 2 from amino acids 26 to 341.
- 63. (New) A method of identifying an inhibitor of IL-13 binding to the IL-13 receptor which comprises:
  - (a) combining a receptor chain protein with IL-13 or a biologically active fragment thereof, said combination forming a first binding mixture;
  - (b) measuring the amount of binding between the receptor chain protein and the IL-13 or fragment in the first binding mixture;
  - (c) combining a compound with the receptor chain protein and the IL-13 or fragment to form a second binding mixture;
  - (d) measuring the amount of binding in the second binding mixture; and
  - (e) comparing the amount of binding in the first binding mixture with the amount of binding in the second binding mixture;

wherein the compound is capable of inhibiting IL-13 binding to the IL-13 receptor when a decrease in the amount of binding of the second binding mixture occurs, and wherein said receptor chain protein comprises an amino acid sequence selected from the group consisting of:

- (1) the mature sequence of the IL-13 receptor chain protein, IL-13 Rβ;
- (2) the extracellular domain sequence of (1); and
- (3) the intracytoplasmic domain sequence of (1).
- 64. (New) The method of claim 63, wherein said protein comprises the amino acid sequence (1).

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- 65. (New) The method of claim 63, wherein said protein comprises the amino acid sequence (2).
- 66. (New) A method of identifying an inhibitor of IL-13 binding to the IL-13Rβ receptor which comprises:
  - (a) measuring the amount of binding between the receptor chain protein IL-13Rβ and IL-13 or a fragment thereof in a first binding mixture;
  - (b) measuring the amount of said binding in a second binding mixture comprising a compound, the receptor chain protein IL-13Rβ and IL-13 or a fragment thereof, and;
  - (c) comparing said amounts of binding to determine whether said compound modulates the biological activity of IL-13Rβ;

wherein the compound is capable of antagonizing IL-13 binding to IL-13Rβ when a decrease in the amount of binding of the second binding mixture occurs, and wherein said receptor chain protein comprises an amino acid sequence selected from the group consisting of:

- (1) the amino acid sequence of SEQ ID NO: 2;
- (2) the amino acid sequence of SEQ ID NO: 2 from amino acids 23 to 342; and
- (3) the amino acid sequence of SEQ ID NO: 2 from amino acids 365 to 380.
- 67. (New) The method of claim 66, wherein said protein comprises the amino acid sequence of SEQ ID NO: 2.
- **68.** (New) The method of claim 66, wherein said protein comprises the amino acid sequence of SEQ ID NO: 2 from amino acids 23-342.
- **69.** (New) A method of identifying an inhibitor of IL-13 binding to the IL-13Rβ receptor which comprises:
  - (a) measuring the amount of binding between the receptor chain protein IL-13Rβ and IL-13 or a fragment thereof in a first binding mixture;
  - (b) measuring the amount of said binding in a second binding mixture comprising a compound, the receptor chain protein IL-13Rβ and IL-13 or a fragment thereof, and;
  - (c) comparing said amounts of binding to determine whether said compound modulates the biological activity of IL-13Rβ;

wherein the compound is capable of antagonizing IL-13 binding to IL-13Rβ when a decrease in the amount of binding of the second binding mixture occurs, and wherein said receptor chain protein comprises an amino acid sequence selected from the group consisting of:

- (1) the mature sequence of the IL-13 receptor chain protein, IL-13 RB;
- (2) the extracellular domain sequence of (1); and
- (3) the intracytoplasmic domain sequence of (1).
- 70. (New) The method of claim 69, wherein said protein comprises the amino acid sequence (1).
- 71. (New) The method of claim 69, wherein said protein comprises the amino acid sequence (2).

72. (New) An isolated IL-13bc (IL13R $\beta$ ) protein comprising an amino acid sequence selected from the group consisting of:

- (a) the amino acid sequence of SEQ ID NO: 2;
- (b) the amino acid sequence of SEQ ID NO: 2 from amino acids 26 to 341;
- (c) the amino acid sequence of SEQ ID NO: 2 from amino acids 363 to 380; and
- (d) fragments of (a) (c) having the ability to bind IL-13 or a biologically active fragment thereof.

(New) The protein of claim 72, comprising the sequence from amino acid 26 to 341 of SEO ID NO: 2.

74. (New) A pharmaceutical composition comprising a protein of claim 72 and a pharmaceutically acceptable carrier.

75. (New) The protein of claim 72, comprising the amino acid sequence of SEQ ID NO: 2 from amino acids 363 to 380.

76. (New) The protein of claim 72, wherein said amino acid sequence is part of a fusion protein.

(New) The protein of claim 76, comprising an Fc fragment.

78. (New) A protein produced according to a process comprising:

- (a) growing a culture of a host cell in a suitable culture medium; and
- (b) purifying the protein from the culture,

wherein said host cell is transformed with a polynucleotide operably linked to an expression control sequence, and wherein said polynucleotide comprises a nucleotide sequence selected

## El from the group consisting of

- (1) a nucleotide sequence encoding the amino acid sequence of SEQ ID 2;
- (2) a nucleotide sequence encoding the IL-13Rβ binding chain varying from the sequence of the nucleotide sequence specified in (1) as a result of degeneracy of the genetic code;
- (3) a nucleotide sequence capable of hybridizing under stringent conditions to
- (1); and
- (4) an allelic variant of the nucleotide sequence specified in (1).
- (New) The protein of claim 78, wherein said nucleotide sequence is that of (1).
- (New) The protein of claim 78, wherein said nucleotide sequence is that of (2).
- (New) The protein of claim 78, wherein said stringent conditions are 52°C in 5xSSC followed by washing at 52°C in 2xSSC.
- (82. (New) The protein of claim 72, comprising SEQ ID NO: 2.
- (83. (New) An isolated IL-13Rβ protein comprising an amino acid sequence selected from the group consisting of:
  - (a) the amino acid sequence of SEQ ID NO: 2;
  - (b) the amino acid sequence of SEQ ID NO: 2 from amino acids 23 to 342;
  - (c) the amino acid sequence of SEQ ID NO: 2 from amino acids 365 to 380; and
  - (d) fragments of (a)-(c) having the ability to bind IL-13 or a biologically active fragment thereof
- 84 (New) The protein of claim 83, comprising the sequence from amino acid 23 to 342 of SEQ ID NO: 2.
- 85. (New) A pharmaceutical composition comprising a protein of claim 83, and a pharmaceutically acceptable carrier.
- 86. (New) The protein of claim 83, comprising the amino acid sequence of SEQ ID NO: 2 from amino acids 365 to 380.
- 87. (New) An isolated IL-13Rβ protein comprising an amino acid sequence selected from the group consisting of:
  - (a) the mature sequence of the IL-13 receptor chain protein, IL-13Rβ;
  - (b) the extracellular domain of sequence (1); and
  - (c) the intracytoplasmic domain sequence of (1).



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- (d) fragments of (a)-(c) having the ability to bind IL-13 or a biologically active fragment thereof.
- 88. (New) A pharmaceutical composition comprising a protein of claim 87, and a pharmaceutically acceptable carrier.
- 89. (New) A method of inhibiting binding of IL-13 to the IL-13Rβ receptor in a mammalian subject, said method comprising administering a therapeutically effective amount of a pharmaceutical composition comprising a protein and a pharmaceutically acceptable carrier, wherein said protein comprises an amino acid sequence selected from the group consisting of:
  - (a) the amino acid sequence of SEQ ID NO: 2;
  - (b) the amino acid sequence of SEQ ID NO: 2 from amino acids 26 to 341; and
  - (c) the amino acid sequence of SEQ ID NO: 2 from amino acids 363 to 380.
- 90. (New) The method of claim 88, wherein said receptor chain protein comprises the amino acid sequence of SEQ ID NO: 2.
- 91. (New) The method of claim 88, wherein said receptor chain protein comprises the amino acid sequence of SEQ ID NO: 2 from amino acids 26 to 341.
- 92. (New) A method of treating an Ig-mediated condition in a mammalian subject, said method comprising administering a therapeutically effective amount of a pharmaceutical composition comprising a protein and a pharmaceutically acceptable carrier, wherein said protein comprises an amino acid sequence selected from the group consisting of:
  - (a) the amino acid sequence of SEQ ID NO: 2;
  - (b) the amino acid sequence of SEQ ID NO: 2 from amino acids 26 to 341;
  - (c) the amino acid sequence of SEQ ID NO: 2 from amino acids 363 to 380; and
  - (d) fragments of (a)-(d) having the ability to bind IL-13 or a biologically active fragment thereof.
- 93. (New) The method of claim 92, wherein said condition is an IgE-mediated condition.
- 94. (New) The method of claim 93, wherein said condition is selected from the group consisting of an allergic condition, asthma and an immune complex disease.
- 95. (New) The method of claim 94, wherein said condition is selected from the group consisting of lupus, nephritis, thyroiditis and Grave's disease.
- 96. (New) The method of claim 92, wherein said protein comprises the amino acid sequence of SEQ ID NO: 2.

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- 97. (New) The method of claim 92, wherein said protein comprises the amino acid sequence of SEQ ID NO: 2 from amino acids 26 to 341.
- 98. (New) The method of claim 91, wherein said protein comprises the amino acid sequence of SEQ ID NO: 2 from amino acids 363 to 380.
- 99. (New) A method of claim 92, wherein said condition is allergy or an inflammatory disease.
- 100. (New) A method of inhibiting binding of IL-13 to the IL-13 receptor in a mammalian subject, said method comprising administering a therapeutically effective amount of a pharmaceutical composition comprising a protein and a pharmaceutically acceptable carrier, wherein said protein comprises an amino acid sequence selected from the group consisting of:
  - (a) the mature sequence of the IL-13 receptor chain protein, IL-13Rβ;
  - (b) the extracellular domain of sequence (1); and;
  - (c) the intracytoplasmic domain sequence of (1).
  - (d) fragments of (a)-(c) having the ability to bind IL-13 or a biologically active fragment thereof.
- 101. (New) A method of treating an Ig-mediated condition in a mammalian subject, said method comprising administering a therapeutically effective amount of a pharmaceutical composition comprising a protein and a pharmaceutically acceptable carrier, wherein said protein comprises an amino acid sequence selected from the group consisting of:
  - (a) the mature sequence of the IL-13 receptor chain protein, IL-13Rβ;
  - (b) the extracellular domain of sequence (1); and;
  - (c) the intracytoplasmic domain sequence of (1).
  - (d) fragments of (a)-(c) having the ability to bind IL-13 or a biologically active fragment thereof.
- 102. (New) The method of claim 101, wherein said condition is an IgE-mediated condition.
- 103. (New) The method of claim 101, wherein said condition is allergy or an inflammatory disease.
- 104. (New) A method of inhibiting binding of IL-13 to the IL-13 receptor in a mammalian subject, said method comprising administering a therapeutically effective amount of a pharmaceutical composition comprising a protein and a pharmaceutically acceptable carrier,

wherein said protein comprises an amino acid sequence selected from the group consisting of:

- (a) the amino acid sequence of SEQ ID NO: 2;
- (b) the amino acid sequence of SEQ ID NO: 2 from amino acids 23 to 342; and
- (c) the amino acid sequence of SEQ ID NO: 2 from amino acids 365 to 380.
- 105. (New) The method of claim 104, wherein said receptor chain protein comprises the amino acid sequence of SEQ ID NO: 2 from amino acids 23 to 342.
- 106. (New) A method of treating an Ig-mediated condition in a mammalian subject, said method comprising administering a therapeutically effective amount of a pharmaceutical composition comprising a protein and a pharmaceutically acceptable carrier, wherein said protein comprises an amino acid sequence selected from the group consisting of:
  - (a) the amino acid sequence of SEQ ID NO: 2;
  - (b) the amino acid sequence of SEQ ID NO: 2 from amino acids 23-342;
  - (c) the amino acid sequence of SEQ ID NO: 2 from amino acids 365 to 380; and
  - (d) fragments of (a)-(c) having the ability to bind IL-13 or a biologically active fragment thereof.
- 107. (New) The method of claim 106, wherein said condition is an IgE mediated condition.
- 108. (New) The method of claim 106, wherein said protein comprises the amino acid sequence of SEQ ID NO: 2 from amino acids 23 to 342.
- 109. (New) The method of claim 106, wherein said protein comprises the amino acid sequence of SEQ ID NO: 2 from amino acids 365 to 380.
- 110. (New) The method of claim 106, wherein said condition is allergy or an inflammatory disease.
- (111) (New) A protein produced according to a process comprising:
  - (a) growing a culture of a host cell in a suitable culture medium; and
  - (b) purifying the protein from the culture,

wherein said host cell is transformed with a polynucleotide operably linked to an expression control sequence, and wherein said polynucleotide comprises a nucleotide sequence selected from the group consisting of

(1) the nucleotide sequence of Fig. (2A and 2B) from nucleotide 53 to nucleotide 1192;

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- (2) a nucleotide sequence encoding the IL-13Rß binding chain varying from the sequence of the nucleotide sequence specified in (1) as a result of degeneracy of the genetic code;
- (3) a nucleotide sequence capable of hybridizing under stringent conditions to (1); and
- (4) an allelic variant of the nucleotide sequence specified in (1).
- 112. (New) The protein of claim 111, wherein said nucleotide sequence comprises that of (1).
- 113. (New) The protein of claim 111 wherein said nucleotide sequence comprises that of (2).

114. (New) The protein of claim 111, wherein said stringent conditions are 52°C in 5xSSC followed by washing at 52°C in 2xSSC.--